GLOBAL STABILITY OF VIRUS PERSISTENCE OF A MATHEMATICAL MODEL OF THE DYNAMICS OF EBOLA VIRUS INFECTION IN HUMAN CELL POPULATION

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Abstract

In this paper, we formulated a mathematical model of the dynamics of Ebola virus infection in human cell population, using ordinary differential equation. We have shown the positivity of solutions of the model equations. The virus free and virus persistence equilibrium points were obtained. We computed the basic reproduction number of the system and shown the global stability of virus persistence equilibrium state.

Keywords: Mathematical model; Ebola virus; Immune response; Positivity; Global stability

1.0 Introduction

Ebola virus is a long filamentous, enveloped virus, consisting of a negative sense single-structural Ribonucleic acid (ssRNA) virus, un-segmented and the genome of the Ebola virus is encoded with Seven structural proteins and two non-structural proteins, such as, Glycoprotein (GP); Nucleoprotein (NP); viral protein 24 (VP24); viral protein 30 (VP30); viral protein 35 (VP35); viral protein 40 (VP40); and RNA-polymerase protein (L-polymerase) [1,2]. Ebola virus belongs to a family of filoviridae [2]. The virus is classified in to five distinct species namely, Reston Ebola virus; Cotes d'Ivoire Ebola virus; Bundibugyo Ebola virus; Sudan Ebola virus; and Zaire Ebola virus [2].

The virus was first discovered in 1976 in Democratic Republic of Congo [3]. Recently, the virus again resurfaced in some West Africa Countries like Guinea, Sierra Leone, Liberia and Nigeria. The latest eruption of the virus was the deadliest in its history [4]. There was an outbreak of the virus between December 2013 and 2015, which was referred to as Zaire Ebola species and known as Ebola virus [2]. It causes serious incidence of haemorrhagic fever and illness in humans who are exposed to the virus [5]. It was reported that at least a total of 23,014 cases occurred of which 9, 840 resulted to deaths [6]. In Nigeria, it was confirmed that a total of 19 cases occurred, of which seven (7) died and twelve (12) survived a fatality rate of 40% compared to 70% fatality rate reported in other Africa Countries [7].

The incubation period of Ebola virus ranges from 2 to 21 days and infectious periods ranges from 4 to 10 days [8]. Meanwhile, it takes an approximation of 31 days to quarantine a patient under investigation of the Ebola virus. Consequently, the symptoms of the Ebola virus are characterized by headaches, fever, vomiting, bleeding diarrhea, and rash [9] in infected person, severe bleeding and shock are usually followed by death. The World Health Organisation has not yet approved any vaccine or antiviral drugs for Ebola virus, rather a trial vaccine is been administer on patients [2]. The spread of the virus and eventual death of infected patients was largely contained (reduced) through early detection and effective contact tracing [1].

Ebola virus is known to cause damage to large variety of cell types including monocytes, macrophages, dendritic cells, endothelial cells, fibroblasts, hepatocytes, and several types of epithelial cells. The primary targets of the virus are dendritic, monocytes and macrophage cells. These are the cells infected in the early stage of the virus infection in the body which spread through the organs [10].

Macrophage encounter APC and release a protein called interlenkin-1(IL-1) that acts as a chemical alarm signal; Helper T-cells respond to interlekin-1 and release interlenkin-2(IL-2) by simultaneously initiating two parallel lines of immune system defence: the cell-mediated response carried out by T-cells, and humoral response carried out by B-cells [11]. The organization of this paper is as follows, in the next section, we present the formation of our model equations, we establish virus free equilibrium state and virus persistence equilibrium and we obtain the basic reproduction number of the infection and determined the global stability of the virus persistence equilibrium state.

2.0 MATERIAL AND METHOD

Model Fermulation

The model equations are formulated using ordinary differential equation. A mathematical model of the dynamics of Ebola virus infection in the cell population and immune response is developed by incorporating humoral immunity, vaccination and clearance of the free virus. The cell population is partitioned into five compartments, these are: U(t) = the Uninfected cells, this is the class in which cell population are free from the virus but open to infection as they interact with the free virus class; I(t) = Infected cells, this the class in which the cell is infected and are infectious; V(t) = free Virus class, this is the class that interact with healthy cells to become infected; T(t) = Cytotoxic T-Lymphocyte (CTL), this is the class that kill infected cell from the cell population; P(t) = Humoral immunity produce by P(t) = CTL), this is the class that clear the free virus from the system.

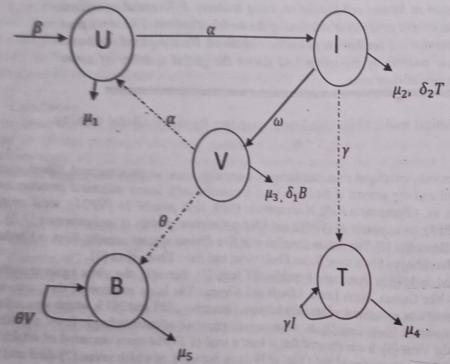


Figure 1: Schematic representation for the model

$$\frac{dU}{dz} = \beta - \mu_1 U - \alpha U \cdot V$$

$$\frac{dI}{dz} = \alpha U \cdot V - \mu_2 I - \delta_1 L T$$

$$\frac{dV}{dz} = \omega I - \mu_3 V - \delta_2 B \cdot V$$

$$\frac{dT}{dz} = \gamma I \cdot T - \mu_4 T$$
(4)

The state of the s	has et the Media
Tourself.	Personal
*	Reproduction rate of animiscisal cell
4	Tribetion rate of unimbered cell
-	Reproduction rate of Virus
*	Reproduction rate of Cyronics: T cells in response to infected cell
	Reproduction take of antibodies in response to Ebole virus exposure
4	Clearance rate of mixted cells by Connoxic T-cell response
4	Clearance rate of the viruses by antibodies
100	Natural death rate of unmirected cell
N	Doub rate of infected coll
70	Death Decay rate of the virus
-	Death Decay rate of Cyrotoxic T-cell
100	Death Decay rate of B-cell secreted antibodies

RESULTS AND DISCUSSION

POSITIVITY OF SOLITIONS OF THE MODEL EQUATIONS

In this section, we to prove that all the solutions to the system of ODEs are positive for all time. 120

Theorem 1: All the solutions of the model equations (1)- (5) are positive for all time t > 0 provided that the initial conditions are positive, that is, U(0) > 0, U(0) > 0, V(0) > 0, T(0) > 0. Then, the solution set U(t), U(t), 172), T(2), B(2) will be positive in Rd

Proof: Assuming that all the parameters are nonnegative, from (1), we have

$$\frac{dV}{dt} = f - f_{1} - aV \cdot V$$

$$\frac{dV}{dt} = f - f_{2} - aV \cdot V$$

$$\frac{dV}{dt} = -\int (f_{2} + aV) dt$$

$$V(t) \ge V(0)e^{-f_{2}t - a}/V(t)dt$$

$$V(t) \ge 0$$

Similarly, we have $I(t) \ge 0$, $V(t) \ge 0$, $T(t) \ge 0$, $B(t) \ge 0$ Thus, for all $t \in [0, \infty)$, U(t), I(t), V(t), I'(t), B(t) will be positive and remain in \mathbb{R}^2 .

EXISTENCE OF EQUILIBRIUM STATE, (F')

Equilibrium States of the Model

At equilibrium states,
$$\frac{dV}{dt} = \frac{dV}{dt} = \frac{dV}{dt} = \frac{dV}{dt} = \frac{dB}{dt} = 0$$

Let $(U, 1, V, T, B) = (U^*, 1^*, V^*, T^*B^*)$

We have the following
$$\beta - \mu_1 U^* - \alpha U^*, V^* = 0$$

$$\alpha U^*, V^* - \mu_2 I^* - \delta_1 I^*, T^* = 0$$

$$\alpha I^* - \mu_3 V^* - \delta_2 B^*, V^* = 0$$

$$\gamma I^*, T^* - \mu_4 T^* = 0$$

$$\theta B^*, V^* - \mu_5 B^* = 0$$
From (12), we have
$$V^* = \frac{\alpha V^*}{\mu_5 + \delta_7 B^*}$$
Substitute V^* in (5) into (11), we have
$$\alpha U^*, \left(\frac{\alpha V^*}{\mu_5 + \delta_5 B^*}\right) - \mu_2 I^* - \delta_1 I^*, T^* = 0$$
(15)

From (16) we have

1 = 0

And
$$\frac{\alpha \omega U^*}{\mu_1 + \delta_2 B^*} - \mu_2 - \delta_1 T^* = 0$$
 (17)

From (13), we have

$$T^*(\gamma I^* - \mu_4) = 0$$

Implies,
$$T^* = 0$$

Or
$$I^* = \frac{\mu_4}{a}$$
 (18)

From (14), we have

$$B^*(\theta.V^*-\mu_5)=0$$

Implies,
$$B^* = 0$$

$$Or V^* = \frac{\mu_5}{\rho} (19)$$

Substitute
$$I^*$$
 in (17), T^* in (18) and B^* in (19) into (15), we have

$$V^* = 0 \tag{20}$$

Substitute I^* in (17) and V^* in (20) into (10), we have

$$U^* = \frac{\beta}{u_*} \tag{21}$$

Therefore, the virus free equilibrium is given as,

$$(U^*, I^*, V^*, T^* B^*) = (U^0, I^0, V^0, T^0 B^0) = \left(\frac{\beta}{\mu_1}, 0, 0, 0, 0, 0\right)$$
 (22)

Substitute $V^* = \frac{\mu_5}{a}$ in (19) into (10), we have

$$\beta - \mu_1 U^* - \alpha U^*(\frac{\mu_5}{\theta}) = 0$$

$$\beta - \mu_1 U^* - \alpha U^* (\frac{\mu_5}{\theta}) = 0$$
Implies
$$U^* = \frac{\beta \theta}{\theta \mu_1 + \alpha \mu_5}$$
Substitute I^* in (18), V^* in (19) and U^* in (23) into (11), we have

$$\alpha \cdot \frac{\beta \theta}{\theta \mu_1 + \alpha \mu_5} \cdot \frac{\mu_5}{\theta} - \mu_2 \cdot \frac{\mu_4}{\gamma} - \delta_1 \frac{\mu_4}{\gamma} \cdot T^* = 0$$
(24)

Simplification of the above gives

$$\frac{\alpha\beta\mu_5}{\theta\mu_1+\alpha\mu_5} - \frac{\mu_2\mu_4}{\gamma} - \frac{\delta_1\mu_4}{\gamma} \cdot T^* = 0$$

Implies,

$$T^* = \frac{\gamma(\alpha\beta\mu_5\gamma - \mu_2\mu_4(\theta\mu_1 + \alpha\mu_5))}{\gamma\delta_1\mu_4(\theta\mu_1 + \alpha\mu_5)}$$

Therefore, we have

Therefore,
$$T^* = \frac{(\alpha\beta\mu_5\gamma - \mu_2\mu_4(\theta\mu_1 + \alpha\mu_5))}{\delta_1\mu_4(\theta\mu_1 + \alpha\mu_5)}$$

Substitute $I^* = \frac{\mu_4}{\gamma}$ in (18) and $V^* = \frac{\mu_5}{\theta}$ in (19) into (12), we have

$$\omega\left(\frac{\mu_4}{\gamma}\right) - \mu_3\left(\frac{\mu_5}{\theta}\right) - \delta_2 B^*.\left(\frac{\mu_5}{\theta}\right) = 0$$

$$\frac{\omega\mu_4}{\gamma} - \frac{\mu_3\mu_5}{\theta} - \frac{\delta_2 B^*\mu_5}{\theta} = 0 \tag{26}$$

From (26), we have

$$B^* = \frac{\omega \mu_4 \theta - \mu_3 \mu_5 \gamma}{\delta_2 \mu_5 \gamma} \tag{27}$$

Therefore, the Virus Persistence Equilibrium (VPE) is given as,

$$(U^*, I^*, V^*, T^* B^*) = (U^{**}, I^{**}, V^{**}, T^{**} B^{**})$$

$$= (\frac{\beta \theta}{\theta \mu_1 + \alpha \mu_5}, \frac{\mu_4}{\gamma}, \frac{\mu_5}{\theta}, \frac{(\alpha \beta \mu_5 \gamma - \mu_2 \mu_4 (\theta \mu_1 + \alpha \mu_5)}{\delta_1 \mu_4 (\theta \mu_1 + \alpha \mu_5)}, \frac{\omega \mu_4 \theta - \mu_3 \mu_5 \gamma}{\delta_2 \mu_5 \gamma})$$
(28)

COMPUTATION OF BASIC REPRODUCTION NUMBER, Ro

The Basic Reproduction number denoted by R_0 is the average number of secondary cases generated by a virus w introduced into a cell population. To control the virus population, it is required that $R_0 < 1$ [12]. In this case infected cell produces on average less than one newly infected case over the course of its infection period. In this the virus will be eliminated from the system and individual will recover from the infection. On the other hand, if & 1 then the infected individual cells produces on average more than one new infected case and the infection will per in the cells population.

We applying next generation matrix operator to compute the basic reproduction number [13]. The basic reproduction number is obtained by dividing the whole population into n compartments in which there are in infected compartment m < n. Let x_i , i = 1,2,3,...,m be the number of infected individual in the i^{th} compartment at time t.

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$$\mathcal{E}_{i}^{\gamma-1} = \begin{bmatrix} \partial F_{i}(K^{0}) \\ \partial S_{i} \end{bmatrix} \begin{bmatrix} \partial V_{i}(K^{0}) \\ \partial S_{i} \end{bmatrix}^{-1}$$

where F_i is the rate of appearance of new infection in compartment I, V_i is the transfer of infection from one compartment I to another and E^0 is the Disease-Free Equilibrium. Using this technique, we have spectral radius (ρ) of next generation matrix, FV^{-1} that is, $R_0 = \rho(FV^{-1})$. Therefore, we have

$$I = \begin{pmatrix} I_1 \\ I_2 \end{pmatrix} = \begin{pmatrix} \omega I \\ \alpha U V \end{pmatrix} \tag{29}$$

pufferentiate (29) partially with respect to time gives

$$F = \begin{pmatrix} \frac{\partial f_1}{\partial V} & \frac{\partial f_1}{\partial I} \\ \frac{\partial f_2}{\partial V} & \frac{\partial f_3}{\partial I} \end{pmatrix} = \begin{pmatrix} 0 & \omega \\ \alpha U & 0 \end{pmatrix}$$
(30)

... virus-free equilibrium (VFE)

$$F = \begin{pmatrix} 0 & \omega \\ \frac{\alpha \beta}{\mu_1} & 0 \end{pmatrix} \tag{31}$$

Given that,
$$v = \begin{pmatrix} v_1 \\ v_2 \end{pmatrix} = \begin{pmatrix} (\mu_2 + \delta_2 B)V \\ (\mu_2 + \delta_1 T)I \end{pmatrix}$$
(32)

oferentiate (32) partially with respect to time, we have,

$$V = \begin{pmatrix} \frac{\partial v_1}{\partial V} & \frac{\partial v_1}{\partial I} \\ \frac{\partial v_2}{\partial V} & \frac{\partial v_2}{\partial I} \end{pmatrix}$$

Implies,

$$V = \begin{pmatrix} (\mu_3 + \delta_2 B^0) & 0 \\ 0 & (\mu_2 + \delta_1 T^0) \end{pmatrix}$$
(33)

$$k_1 = (\mu_3 + \delta_2 B^0) k_1 = (\mu_2 + \delta_1 T^0)$$
(34)

We have,
$$V^{-1} = \frac{adjv}{adjv}$$
(35)

Implies,

$$V^{-1} = \frac{aa_1V}{\det V} = \frac{1}{k_1k_2} \begin{pmatrix} k_2 & 0 \\ 0 & k_1 \end{pmatrix}$$

$$V^{-1} = \frac{1}{\det V} \begin{pmatrix} k_2 & 0 \\ 0 & k_1 \end{pmatrix}$$
(16)

Multiplying (31) and (36) gives

$$FV^{-1} = \frac{1}{k_1 k_2} \begin{pmatrix} 0 & \omega \\ \frac{\alpha \beta}{\mu_*} & 0 \end{pmatrix} \begin{pmatrix} k_2 & 0 \\ 0 & k_1 \end{pmatrix}$$

$$\tag{37}$$

solving (37), we have

$$FV^{-1} = \frac{1}{k_1 k_2} \begin{pmatrix} 0 + 0 & \omega k_1 \\ \frac{\alpha \beta k_2}{\mu_1} - 0 & 0 + 0 \end{pmatrix}$$

$$FV^{-1} = \begin{pmatrix} 0 & \frac{\omega k_1}{k_1 k_2} \\ \frac{\alpha \beta k_2}{k_1 k_2 \mu_1} & 0 \end{pmatrix}$$

Implies,

$$FV^{-1} = \begin{pmatrix} 0 & \frac{\omega}{k_2} \\ \frac{\alpha\beta}{k_1\mu_1} & 0 \end{pmatrix} \tag{3B}$$

The characteristics equation of (38) gives

$$\begin{vmatrix} -\lambda & \frac{\omega}{k_1} \\ \frac{a\beta}{k_2 \mu_1} & -\lambda \end{vmatrix} = 0$$

$$\lambda^2 - \left(\frac{\omega}{k_1}, \frac{a\beta}{k_2 \mu_1}\right) = 0$$

$$\lambda^2 = \left(\frac{\omega}{k_2}, \frac{a\beta}{k_2 \mu_2}\right) = \frac{a\omega\beta}{k_1 k_2 \mu_2}$$
Implies.
$$\lambda^2 = \frac{a\omega\beta}{(\mu_1 + \delta_2 T^0)(\mu_2 + \delta_2 S^0)\mu_2}$$

$$\lambda = \frac{a\omega\beta}{(\mu_1 + \delta_2 T^0)(\mu_2 + \delta_2 S^0)\mu_2}$$

$$\lambda \text{ is the spectral radius of } \rho(FV^{-1})$$

$$R_3 = \sqrt{\frac{a\omega\beta}{(\mu_1 + \delta_2 T^0)(\mu_2 + \delta_2 S^0)\mu_2}}$$
(41)

The equation (41) is the basic reproduction number of infected cell of system (1)-(5).

GLOBAL STABILITY OF VIRUS PERSISTENCE EQUILIBRIUM STATE (E")

Luapunov function of the Goh-Voltera type which is nonlinear is used for proving of the Global asymptotic stability of the Endemic Equilibrium Point (EEP) which is known as Virus Persistence and Immunes Persistence Equilibrium State, (VIEP)

Theorem 2: The unique virus and immunes persistence equilibrium of the model (1)-(5), is globally asymptotically stable (GAS) if R, > 1 & R, > 1

Consider the model (1)+5), let $R_a > 1$. Consider the following non-linear Lyapunov function for the sub-system model involving the state variables, U, V, and I. Let

$$F = U - U^{-n} - U^{-n} \ln(\frac{u}{u^{-n}}) + V - V^{-n} - V^{-n} \ln(\frac{v}{u^{-n}}) + A(I - I^{-n} - I^{-n} \ln(\frac{t}{u^{-n}}))$$
(42)

The Lyapunov derivative from equation (42), F. gives

$$\ddot{F} = (\dot{U} - \frac{U''}{U}\dot{U}) + (\dot{V} - \frac{V''}{U}\dot{V}) + A(I - \frac{V''}{U}\dot{I})$$
(43)

Substitute the night hand equations in the model (1)-(3) into the expression in (43), we have

$$F = \beta - \mu_{2}U - \alpha U \cdot V - \frac{u^{-}}{u}(\beta - \mu_{2}U - \alpha U \cdot V) + \omega l - \mu_{3}V - \delta_{2}\beta \cdot V - \frac{v^{-}}{V}(\omega l - \mu_{3}V - \delta_{2}\beta \cdot V) + A(\alpha U \cdot V - \mu_{2}I - \delta_{1}l \cdot T - \frac{v^{-}}{I}(\alpha U \cdot V - \mu_{2}I - \delta_{1}l \cdot T))$$
(44)

Simplification of (44) given

$$\ddot{F} = \beta - \mu_{1}U - \alpha U.V - \frac{v^{**}\beta}{v} + \mu_{1}U^{***} + \alpha U^{***}.V
+ \omega I - \mu_{3}V - \delta_{2}B.V - \frac{v^{**}}{v}\omega I + \mu_{3}V^{***} + \delta_{2}B.V^{***}
+ A(\alpha U.V - \mu_{2}I - \delta_{1}I.T - \frac{v^{**}}{v}\alpha U.V + \mu_{2}I^{***} + \delta_{1}I^{***}.T)$$
(45)

Substitute the equation (6) of the model into equation (45)

$$\dot{F} = \beta - \mu_1 U - \alpha U \cdot V - \frac{u^{-\alpha}\beta}{u} + \mu_1 U^{-\alpha} + \alpha U^{-\alpha} \cdot V
+ \omega I - \mu_3 V - \delta_2 \beta \cdot V - \frac{v^{-\alpha}}{v} \omega I + \mu_3 V^{-\alpha} + \delta_2 \beta \cdot V^{-\alpha}
+ \alpha A U \cdot V - \mu_2 A I - \delta_1 A I \cdot T - \frac{r^{-\alpha}}{u} \alpha A U \cdot V + \mu_2 A I^{-\alpha} + \delta_1 A I^{-\alpha} \cdot T$$
(46)

At endemic steady state, we note that from equation (1)-(5), we have the following relation

$$\beta = \mu_{2}U^{**} + \alpha U^{**}V^{**}$$

$$\omega U^{**} = \mu_{2}V^{**} + \delta_{2}B^{**}.V^{**}$$

$$\omega U^{**} = \mu_{2}U^{**} + \delta_{3}U^{**}.T^{**}$$

$$(47)$$

$$\omega U^{**}.V^{**} = \mu_{2}U^{**} + \delta_{3}U^{**}.T^{**}$$

$$(49)$$

$$W^{**}.T^{**} = \mu_{4}T^{**}$$

$$(50)$$

$$\Theta B^{**}.V^{**} = \mu_{5}B^{**}$$

$$(51)$$

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(At) normalis the expression of it in equation (47) most be equation (44)

$$+ mAO \cdot F = m_0AF - d_0AI \cdot F = \frac{1}{2} mAO \cdot F + m_0AI'' + d_0AI'' \cdot F$$
 (87)

We noted the feature, where the inflected classes are without the three stars on them, (i.e V, I). these terms are now collected regether, we have

We openie opinion (33) to sero (3), and find the value of A, now to determine A, collecting play forms in inflicted (1) only, or the coefficients of the infected term from equation (53), we have

$$ml = \mu_0 Al = \delta_1 Al, F = 0 \tag{51}$$

bughles.

$$\alpha \dot{\beta} = A(\rho_0 + \delta_1, T)\dot{\beta} = 0 \tag{54}$$

Making A subject of formula, we have,

$$A = \frac{n}{(m_1 + d_1 + 1)} \tag{55}$$

to continu that the expression in (33) is zero (0), we have

$$-(aU)V + aU^{**}(V) + ad - \mu_3 V - \delta_2 B.V + (aU)A.V + (-\mu_2 I - \delta_1 I.T)A = 0$$
 (56)

HAMMA

$$-\left(\frac{n_2+\delta_1T}{V}\right)tV + \alpha tV^{**}\left(\frac{n_2+\delta_1T}{\alpha V^{**}}\right)t + \omega t - \omega t + \left(\frac{n_2+\delta_1T}{V}\right)t\left(\frac{\omega}{(\mu_2+\delta_1T)}\right)V - \left(\frac{\omega}{n_2}t + \delta_1t, T\right)\left(\frac{\omega}{(\mu_2+\delta_1T)}\right) = 0$$
(57)

Kerkerkerk

$$-(\kappa_2 + \delta_1 T)t + (\kappa_2 + \delta_1 T)t + \omega t - \omega t + (t)(\omega) - (t)(\omega) = 0$$
 (58)

Since (53) is equal to zero (0), the expression of F in (52), reduced to the following.

Since (33) is equal to zero (0), the expression of
$$F$$
 in (52), reduced to the following,
$$F = \mu_1 U^{**} + \alpha U^{**}V^{**} - \mu_1 U - \frac{\mu_1 U^{**}}{V} - \frac{\alpha U^{**}V^{**}}{V} + \mu_1 U^{**} - \frac{\nu^{**}}{V} \omega I + \mu_3 V^{**} + \delta_2 B. V^{**} - \frac{\nu^{**}}{I} \alpha \left(\frac{\omega}{(\mu_2 + \delta_1.T)}\right) U.V + \frac{\omega}{I} \left(\frac{\omega}{(\mu_2 + \delta_1.T)}\right) I^{**} + \delta_1 \left(\frac{\omega}{(\mu_2 + \delta_1.T)}\right) I^{**}. T$$
(59)

Since $\frac{1}{\mu_2 + \delta_1.T} = \frac{\omega}{\alpha U^{**}V^{**}}$ and $\frac{\omega}{(\mu_3 + \delta_2.B)} = \frac{\nu^{**}}{I^{**}}$ at steady state from equation (2) and (3), then we substitute into equation (34), we have.

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{v}{v^{**}} - \frac{v^{**}}{v} \right) + \alpha U^{**} V^{**} - \frac{\alpha v^{**2} v^{**}}{v} - \frac{v^{**}}{v} \omega I + \omega I^{**} - \frac{t^{**} \omega}{\alpha v^{**} v^{**}} \right) U \cdot V + \mu_2 \left(\frac{t^{**} \omega}{\alpha v^{**} v^{**}} \right) I^{**} + \delta_1 \left(\frac{\omega I^{**}}{\alpha v^{**} v^{**}} \right) I^{**} \cdot T$$

$$(60)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{V^{**}}{U} \right) + \alpha U^{**} V^{**} \left(1 - \frac{V^{**}}{U} \right) + \omega I^{**} \left(1 - \frac{V^{**}}{V^{**}} \right)$$

$$- \frac{I^{**}}{I} \left(\frac{I^{**} \omega U V}{U^{**} V^{**}} \right) + \left(\alpha U^{**} V^{**} \right) \left(\frac{\omega I^{**}}{\alpha U^{**} V^{**}} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{V^{**}}{U} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U} \right) + \omega I^{**} \left(1 - \frac{V^{**}}{V^{I*}} \right)$$

$$- \frac{I^{**}}{I} \left(\frac{I^{**} \omega U V}{U^{**} V^{**}} \right) + (\omega I^{**})$$

$$- \frac{I^{**}}{I} \left(\frac{I^{**} \omega U V}{U^{**} V^{**}} \right) + \omega I^{**} \left(1 - \frac{V^{**}}{V^{I*}} \right)$$

$$+ \omega I^{**} \left(1 - \left(\frac{I^{**} U V}{U^{**} V^{**}} \right) \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{V^{**}}{V^{I*}} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{V^{**}}{V^{I*}} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{V^{**}}{V^{I*}} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{V^{**}}{U^{**}} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U^{**}} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{V^{**}}{U^{**}} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U^{**}} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U^{**}} \right) + \omega I^{**} \left(1 - \frac{U^{**}}{U^{**}} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U^{**}} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U^{**}} \right) + \omega I^{**} \left(1 - \frac{U^{**}}{U^{**}} \right) + \alpha U^{**} V^{**} \left(1 - \frac{U^{**}}{U^{**}} \right)$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U}{U^{**}} \right) + \omega I^{**} \left(2 - \frac{U}{U^{**}} - \frac{U}{U^{**}} \right) + \omega I^{**} \left(2 - \frac{U}{U^{**}} \right) + \omega I^{**} \left($$

$$-\frac{I^{**}}{I}\left(\frac{I^{**}\omega UV}{U^{**}V^{**}}\right) + (\omega I^{**}) \tag{62}$$

$$+\omega l^{**}\left(1-\left(\frac{l^{**}uv}{n^{**}v^{**}}\right)+\right) \tag{63}$$

$$\dot{F} = \mu_1 U^{**} \left(2 - \frac{U}{U^{**}} - \frac{U^{**}}{U} \right) + \omega I^{**} \left(2 - \frac{V^{**}}{VI^{**}} I - \frac{I^{**}UV}{U^{**}VI^{**}} \right) + \alpha U^{**}V^{**} \left(1 - \frac{U^{**}}{U} \right)$$
(64)

Finally, since the arithmetic mean exceeds the geometric mean, i. e
$$AM = \frac{x_1 + x_2 + x_3 + \dots + x_n}{n} \ge \sqrt[n]{x_1 + x_2 + x_3 + \dots + x_n} = GM [14]$$

It is follows that.

$$\mu_1 U^{\bullet \bullet} \left(2 - \frac{v}{v^{\bullet \bullet}} - \frac{v^{\bullet \bullet}}{v} \right) \le 0$$

$$\omega I^{\bullet \bullet} \left(2 - \frac{v^{\bullet \bullet}}{v_I^{\bullet \bullet}} - \frac{i^{\bullet \bullet} v v}{v^{\bullet \bullet} v^{\bullet}} \right) \le 0$$

$$\alpha U^{**}V^{**}\left(1 - \frac{U^{**}}{U}\right) \le 0 \tag{65}$$

Furthermore, since all the model parameters are non-negative, it follows that $\dot{F} \leq 0$ for $R_o > 1$, thus \dot{F} is a Lyapunov function for sub system of the model (1)- (5) and is globally asymptotically stable. Therefore, it follows by Lassalle's Invariance Principle [14] that,

$$\lim_{t \to \infty} U(t) = U^{**}, \quad \lim_{t \to \infty} V(t) = V^{**} \quad \text{and} \quad \lim_{t \to \infty} I(t) = I^{**}$$
 (66)

Since $I(t) \to I^*$ as $t \to \infty$, it follows from the equation (4) at steady state, that

$$(-) \to 0 \qquad \to \infty \tag{67}$$

The epidemiological implication of the Theorem I, is that Ebola virus will be establish in the system (or Body) when > 1 and the immune responses against the Ebola virus in the cell population move toward zero as the virus

CONCLUSION

This paper shown the positivity of solutions of the model equations (1)-(5), which implies, that the solutions remain positive for all time, t and the model is epidemiologically meaningful and well pose. Equation (22) and (28) showed the existence of virus free and virus persistence equilibria points. Equation (41) computed the basic reproduction number of the system and Equations (65), (66) & (67) shown the global stability of virus persistence equilibrium state and found that the virus persistence equilibrium is globally asymptotical stable (GAS) when $\square_0 > 1$.

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